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netic mapping; gene expression control; promoter;
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                                                                                              The present sequence is a putative protein fragment from Arabidopsis thaliana. Its coding sequence was isolated by carrying out RT-PCR on all of the mRNA obtained from the plant, and creating a cDNA library which could then be sequenced, allowing the putative protein sequence(s) to be obtained. This sequence may be useful for protein identification and for siding in the elucidation of signal transduction and metabolic pathways. Its coding sequence has a use in the control of gene expression as a promoter; coding sequence, 3'UTR or termination sequence, for controlling the behaviour of a gene within the chromosome, as a cool for use in penaltic mapping, including a use in hybridisation assays, for recognition or isolation of similar DNA fragments, or for the identification of a
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AAG47089 Sequence 1176 AA; \vdash MSITCELLNL TSKKAKKSSS SDKKWLKKPL FFLILCGSLV IVLVMFLRLG Length: 1176 June 21, 2005 13:40 Type: P Check: 2954

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651 601 551 501 451 401 351 301 251 201 151 101 51 SGRLELENVP FDMRFILDNV SSLLSGKANE KGIELAVYVS RSQKEETDSC NGEEKVLYRH TPMNGVLGML KMLMDTDLDA KQMDYAQTAH GSGKDLTSLI NEVLDQAKIE SEIGDISEQH RVEATIGYLG KKMETEDQTV KIPSAIDQRT FEEYTERTNF WDTNEEIIMK RRETLANMCD QQPENLEEEN SPGMWTNYGI TCSLSVRSDK QETRGLPWNL GLGHSISSTS LAPKKQQQRA HEEGLEQGLS SYLRNAWWCL ILGVLVCHKI YVSHSKARGE ISSLDFGDPS ASYDMPSLVE ASGKGVLTSP VQDCVPENFD RNHEMHCRFK HKLPIPWTAI KLLHQLASKQ TIAVDVYDTT FKLLKSNHLG PAPIQDEYAP VIFAQETVSH IVSVDMMSGE ERPLTSGVAY ALKVPHSERE ERARVLODOF NVSLNHVHAL QTSSRGAGRW QNVTRSEIHD LVSLFSDSDQ COMMRELKAR VVLTFAVYDT RKNILLLGIL AEAADIAKSQ SQVPDVVVGD TPSILVLVIT NTSGLIKMYG SILVSTFHHG FLATVSHEIR SLPPDATEEQ KFEKEHGWAI GGVSFSVWWF CMCGNLEPIL VISFECHKES

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